D. N.Sona	CRF Errors Corrected by th STIC Systems Eanch CRF Processing Date: 11/29/2002 CRF Processing Date: 11/29/2002
	mber: 09/69, 220 E Edited by: V rifled by: (STIC staff)
	changed the margins in cases where the sequence text was "wrapped" down to the next line.
	dited a format error in the Current Application Data section, specifically:
E	dited the Current Application Data section with the actual current number. The number inputted by the pplicant was the prior application data; or other
Α	dded the mandatory heading and subheadings for "Current Application Data".
Ε	dited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
С	changed the spelling of a mandatory field (the headings or subheadings), specifically:
C	corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
In	nserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
C a _l	orrected subheading placement. All responses must be on the same line as each subheading. If the pplicant placed a response below the subheading, this was moved to its appropriate place.
ļı	nserted colons after headings/subheadings. Headings edited included:
_	Deleted extra, invalid, headings used by an applicant, specifically:
 I	Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of file; page numbers throughout text; other invalid text, such as
ı	Inserted mandatory headings, specifically:
(Corrected an obvious error in the response, specifically:
	Edited identifiers where upper case is used but lower case is required, or vice versa. TECH CENTER 1600/
(Corrected an error in the Number of Sequences field, specifically:
	A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
D: d:	eleted endIng stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error use to a PatentIn bug). Sequences corrected:
_	Other:
-	
_	the first Office

*Examiner: The abov corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.



1600

RAW SEQUENCE LISTING DATE: 11/29/2002 PATENT APPLICATION: US/09/691,220 TIME: 17:36:40

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4 <110> APPLICANT: WEI, Ming-Hui et al.

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         RECEPTORS, NUCLEIC ACID MOLECULES ENCODING HUMAN NUCLEAR
         HORMONE RECEPTORS, AND USES THEREOF
11 <130> FILE REFERENCE: CL000893
13 <140> CURRENT APPLICATION NUMBER: US 09/691,220
14 <141> CURRENT FILING DATE: 2000-10-19
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76 Glu Ile Val Pro Ser Pro Pro Ser Pro Pro Pro Leu Pro Arg Ile Tyr
77 65
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80 Val Ser Ala Cys Glu Gly Cys Lys Gly Phe Phe Arg Arg Ser Ile Gln
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                                   105
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88 Lys Lys Glu Val Pro Lys Pro Glu Cys Ser Glu Ser Tyr Thr Leu Thr
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90 Pro Glu Val Gly Glu Leu Ile Glu Lys Val Arg Lys Ala His Gln Glu
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94 Ser Glu Gln Arg Val Ser Leu Asp Ile Asp Leu Trp Asp Lys Phe Ser
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Input Set : A:\PTO.AMC.TXT

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Input Set : A:\PTO.AMC.TXT

Output Set: N:\CRF4\11292002\I691220.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:3; N Pos. 15107,15108,15109,15110,15111,15112,15113,15114,15115,15116
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-ahi

1600

RAW SEQUENCE LISTING DATE: 11/22/2002 PATENT APPLICATION: US/09/691,220 TIME: 08:51:00

Input Set : A:\SEQLIST_893.TXT

Output Set: N:\CRF4\11222002\I691220.raw

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- 8 HORMONE RECEPTORS, AND USES THEREOF
- 11 <130> FILE REFERENCE: CL000893
- 13 <140> CURRENT APPLICATION NUMBER: US 09/691,220
- 14 <141> CURRENT FILING DATE: 2000-10-19
- 16 <160> NUMBER OF SEQ ID NOS: 4
- 18 <170> SOFTWARE: FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

425	<21	1> L	EQ II ENGTI YPE:	H: 45						Q.)	_			Do	e No	iskette Neede d Skette Needed
					Rattus norvegicus				A Co			Co) We crow			
			EQUEI					-							\mathcal{P}_{\emptyset}	
430	Met	Tyr	Glu	Ser	Val	Glu	Val	Gly	Gly	Leu	Thr	Pro	Ala	Pro	Asn	Pro
431	1				5					10					15	
432 433	Phe	Leu	Val	Val 20	Asp	Phe	Tyr	Asn	Gln 25	Asn	Arg	Ala	Cys	Leu 30	Leu	Gln
434 435	Glu	Lys	Gly 35	Leu	Pro	Ala	Pro	Gly 40	Pro	Tyr	Ser	Thr	Pro 45	Leu	Arg	Thr
436 437	Pro	Leu 50	Trp	Asn	Gly	Ser	Asn 55	His	Ser	Ile	Glu	Thr 60	Gln	Ser	Ser	Ser
	Ser		Glu	Ile	Val	Pro	Ser	Pro	Pro	Ser	Pro	Pro	Pro	Leu	Pro	Arg
439	65					70					75					80
440 441	Ile	Tyr	Lys	Pro	Cys 85	Phe	Val	Cys	Gln	Asp 90	Lys	Ser	Ser	Gly	Tyr 95	His
442 443	Tyr	Gly	Val	Ser 100	Ala	Суѕ	Glu	Gly	Cys 105	Lys	Gly	Phe	Phe	Arg 110	Arg	Ser
	Ile	Gln	Lys 115	Asn	Met	Val	Tyr	Thr 120	Cys	His	Arg	Asp	Lys 125	Asn	Cys	Ile
446 447	Ile	Asn 130	Lys	Val	Thr	Arg	Asn 135	Arg	Cys	Gln	Tyr	Cys 140	Arg	Leu	Gln	Lys
448	Cys		Glu	Val	Gly	Met	Ser	Lys	Glu	Ser	Val	Arg	Asn	Asp	Arg	Asn
449	145				_	150		_			155					160
450 451	Lys	Lys	Lys	Lys	Glu 165	Thr	Pro	Lys	Pro	Glu 170	Cys	Ser	Glu	Ser	Tyr 175	Thr
452 453	Leu	Thr	Pro	Glu 180	Val	Gly	Glu	Leu	Ile 185	Glu	Lys	Val	Arg	Lys 190	Ala	Asn
454	Gln	Glu	Thr	Phe	Pro	Ala	Leu	Cys	Gln	Leu	Gly	Lys	Tyr	Thr	Thr	Asn.

Input Set : A:\SEQLIST_893.TXT

```
455
                195
                                     200
                                                         205
     456 Asn Ser Ser Glu Gln Arg Val Ser Leu Asp Ile Asp Leu Trp Asp Lys
             210
                                 215
                                                     220
     458 Phe Ser Glu Leu Ser Thr Lys Cys Ile Ile Lys Thr Val Glu Phe Ala
     459 225
                             230
                                                 235
     460 Lys Gln Leu Pro Gly Phe Thr Thr Leu Thr Ile Ala Asp Gln Ile Thr
                         245
                                             250
     462 Leu Leu Lys Ala Ala Cys Leu Asp Ile Leu Ile Leu Arg Ile Cys Thr
                                         265
     464 Arg Tyr Thr Pro Glu Gln Asp Thr Met Thr Phe Ser Asp Gly Leu Thr
                275
                                     280
     466 Leu Asn Arg Thr Gln Met His Asn Ala Gly Phe Gly Pro Leu Thr Asp
                                 295
                                                     300
     468 Leu Val Phe Ala Phe Ala Asn Gln Leu Leu Pro Leu Glu Met Asp Asp
                             310
                                                 315
     470 Ala Glu Thr Gly Leu Leu Ser Ala Ile Cys Leu Ile Cys Gly Asp Arg
                         325
                                             330
     472 Gln Asp Leu Glu Gln Pro Asp Lys Val Asp Met Leu Gln Glu Pro Leu
                     340
                                         345
     474 Leu Glu Ala Leu Lys Val Tyr Val Arg Lys Arg Arg Pro Ser Gln Pro
                                     360
                 355
     476 His Met Phe Pro Lys Met Leu Met Lys Ile Thr Asp Leu Arg Ser Ile
             370
                                 375
     478 Ser Ala Lys Gly Ala Glu Arg Val Ile Thr Leu Lys Met Glu Ile Pro
     479 385
                             390
                                                 395
     480 Gly Ser Met Pro Pro Leu Ile Gln Glu Met Leu Glu Asn Ser Glu Gly
                                             410
     482 Leu Asp Thr Leu Ser Gly Gln Ser Gly Gly Gly Thr Arg Asp Gly Gly
                     420
                                         425
     484 Gly Leu Ala Pro Pro Pro Gly Ser Cys Ser Pro Ser Leu Ser Pro Ser
         435
                                     440
     486 Ser His Arg Ser Ser Pro Ala Thr Gln Ser Pro
     487
            450
                                 455
E--> 489 (1)
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VERIFICATION SUMMARY

DATE: 11/22/2002

PATENT APPLICATION: US/09/691,220

TIME: 08:51:01

Input Set : A:\SEQLIST 893.TXT

Output Set: N:\CRF4\11222002\I691220.raw

L:177 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:2280 L:390 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:15060 L:391 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:15120 L:489 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:4